

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: September 21, 2004, 17:07:10 ; Search time 128 Seconds
(without alignments)

672.373 Million cell updates/sec

Title: US-09-875-456A-14

Perfect score: 1444

Sequence: 1 MGRLLLVGAAALVSSACGG.....QRVVLGPGIIIRCVSRGVV 268

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 1342398 seqs, 32113274 residues

Total number of hits satisfying chosen parameters: 1342398

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	1444	100.0	268	9	US-09-875-456A-14
2	1444	100.0	268	15	US-10-401-916-14
3	783	54.2	218	14	US-10-142-201B-8
4	751	52.0	218	9	US-09-997-579-44
5	749	51.9	218	13	US-10-029-191-20
6	258.5	17.9	159	9	US-09-997-579-22
7	258.5	17.9	215	14	US-09-997-579-2
8	258.5	17.9	215	14	US-10-142-201B-11
9	257.5	17.8	159	9	US-09-997-579-23
10	257.5	17.8	215	13	US-10-029-191-4
11	257.5	17.8	215	9	US-09-997-579-1
12	257.5	17.8	215	13	US-10-029-191-2
13	257.5	17.8	215	14	US-10-142-201B-12
14	251	17.4	111	13	US-10-029-191-5
15	115.5	8.0	209	14	US-10-095-131A-20

Sequence 24, Appli	209	14	US-10-095-131A-24
Sequence 4, Appli	269	14	US-10-095-131A-4
Sequence 6, Appli	269	14	US-10-095-131A-6
Sequence 8, Appli	269	14	US-10-095-131A-8
Sequence 34, Appli	159	14	US-10-095-131A-34
Sequence 38, Appli	159	14	US-10-095-131A-38
Sequence 42, Appli	159	14	US-10-095-131A-42
Sequence 46, Appli	199	14	US-10-095-131A-46
Sequence 18, Appli	199	14	US-10-095-131A-18
Sequence 22, Appli	209	14	US-10-095-131A-22
Sequence 364, App	209	14	US-10-095-131A-22
Sequence 530, App	269	10	US-09-946-374-364
Sequence 530, App	269	12	US-10-147-493-530
Sequence 530, App	269	12	US-10-145-127-530
Sequence 530, App	269	12	US-10-160-503-530
Sequence 530, App	269	12	US-10-143-118-530
Sequence 530, App	269	12	US-10-144-983-530
Sequence 530, App	269	12	US-10-158-787-530
Sequence 530, App	269	12	US-10-081-056-252
Sequence 530, App	269	12	US-10-140-024-530
Sequence 530, App	269	12	US-10-140-808-530
Sequence 364, App	269	12	US-10-006-485A-364
Sequence 364, App	269	12	US-10-013-907A-364
Sequence 364, App	269	12	US-10-015-499A-364
Sequence 364, App	269	12	US-10-013-910A-364
Sequence 364, App	269	12	US-10-152-405-530
Sequence 530, App	269	12	US-10-127-852A-530
Sequence 530, App	269	12	US-10-127-900A-530
Sequence 530, App	269	12	US-10-128-685A-530
Sequence 364, App	269	12	US-10-228-254A-364
Sequence 530, App	269	12	US-10-131-820A-530

ALIGNMENTS

RESULT 1

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US-09-875-456A-14
; Sequence 14, Application US/09875456A
; Patent No. US20020045229A1
; GENERAL INFORMATION:
; APPLICANT: Qin, Ning
; APPLICANT: D'Andrea, Michael
; TITLE OF INVENTION: DNAs encoding human betala sodium channel subunit
; FILE REFERENCE: ORT-1221
; CURRENT APPLICATION NUMBER: US/09/875,456A
; CURRENT FILING DATE: 2001-09-10
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 14
; LENGTH: 268
; TYPE: PRT
; ORGANISM: Homo sapiens
; US-09-875-456A-14
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Query Match	100.0%;	Score 1444;	DB 9;	Length 268;
Best Local Similarity	100.0%;	Pred. No. 4.6e-134;		
Matches 268;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;
Qy	1	MGRLLLVGAAALVSSACGGVEVDSETEAVYGVTFKILCISCKRSETNAETFTWTFR	60	
Db	1	MGRLLLVGAAALVSSACGGVEVDSETEAVYGVTFKILCISCKRSETNAETFTWTFR	60	
Qy	61	QKGTEEFVKILRYENEVLQLEEDERPEGRVWVWNGSRGKLODLISIFITVTVNHSGDYE	120	
Db	61	QKGTEEFVKILRYENEVLQLEEDERPEGRVWVWNGSRGKLODLISIFITVTVNHSGDYE	120	
Qy	121	CHVYRLAFFENYEHTSVVKKIHEVVDKESGAACPFTVTHRRARWRDQAVDRGTGWL	180	
Db	121	CHVYRLAFFENYEHTSVVKKIHEVVDKESGAACPFTVTHRRARWRDQAVDRGTGWL	180	
Qy	181	CAWFANRPPQRAEGSGSPSCPLQLWPLFLSSPRGSGMPVPHRRSGYRTQLCHLCOMTS	240	

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Db 181 CWPANRPOORAEGBGSSPCPLQLWPLFLSSPRGQSMVPVPHRRSGYRTOLCHLCMTS 240
Qy 241 GRCLLSLSORVVLGLPGIIRVCVSRGVV 268
Db 241 GRCLLSLSORVVLGLPGIIRVCVSRGVV 268

RESULT 2
US-10-401-916-14
; Sequence 14, Application US/10401916
; Publication No. US20040002439A1
; GENERAL INFORMATION:
; APPLICANT: Qin, Ning
; APPLICANT: Codd, Ellen
; APPLICANT: D'Andrea, Michael
; TITLE OF INVENTION: DNAs encoding human betala sodium channel subunit
; FILE REFERENCE: ORT-1221
; CURRENT APPLICATION NUMBER: US/10/401,916
; PRIOR FILING DATE: 2003-03-28
; PRIOR APPLICATION NUMBER: US/09/875,456A
; PRIOR FILING DATE: 2001-09-10
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 14
; LENGTH: 268
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-401-916-14

Query Match 100.0%; Score 1444; DB 15; Length 268;
Best Local Similarity 100.0%; Pred. No. 4.6e-134;
Matches 268; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MGRLLALVVGAAVLSVSSACGGCVVDSETEAVYGMTFKILCISCKRSETNAETFTWTFR 60
Db 1 MGRLLALVVGAAVLSVSSACGGCVVDSETEAVYGMTFKILCISCKRSETNAETFTWTFR 60

Qy 61 QKGTTEFVKILRYENEVLQLEDEDERFEGRVVWNGSRGTDQLDLSIFITNTVYHSGDYE 120
Db 61 QKGTTEFVKILRYENEVLQLEDEDERFEGRVVWNGSRGTDQLDLSIFITNTVYHSGDYE 120

Qy 121 CHVYRLFFENYEHNTSVVKKIHIEVVDKESGAA 155
Db 121 CHVYRLFFENYEHNTSVVKKIHIEVVDKESGAA 155

RESULT 3
US-10-142-201B-8
; Sequence 8, Application US/10142201B
; Publication No. US20030022205A1
; GENERAL INFORMATION:
; APPLICANT: Millenium Pharmaceuticals Inc.
; APPLICANT: Curtis, Rory A.J.
; TITLE OF INVENTION: 98359, A SODIUM CHANNEL BETA 4 SUBUNIT,
; FILE REFERENCE: AND USES THEREFOR
; CURRENT APPLICATION NUMBER: US/10/142,201B
; PRIOR FILING DATE: 2002-05-09
; PRIOR APPLICATION NUMBER: US 60/289,893
; PRIOR FILING DATE: 2001-05-09
; NUMBER OF SEQ ID NOS: 12
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8
; LENGTH: 218
; TYPE: PRT

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; ORGANISM: Homo Sapiens
US-10-142-201B-8

Query Match 54.2%; Score 783; DB 14; Length 218;
Best Local Similarity 96.8%; Pred. No. 7.1e-69;
Matches 150; Conservative 0; Mismatches 5; Indels 0; Gaps 0;

Qy 1 MGRLLALVVGAAVLSVSSACGGCVVDSETEAVYGMTFKILCISCKRSETNAETFTWTFR 60
Db 1 MGRLLALVVGAAVLSVSSACGGCVVDSETEAVYGMTFKILCISCKRSETNAETFTWTFR 60

Qy 61 QKGTTEFVKILRYENEVLQLEDEDERFEGRVVWNGSRGTDQLDLSIFITNTVYHSGDYE 120
Db 61 QKGTTEFVKILRYENEVLQLEDEDERFEGRVVWNGSRGTDQLDLSIFITNTVYHSGDYE 120

Qy 121 CHVYRLFFENYEHNTSVVKKIHIEVVDKESGAA 155
Db 121 CHVYRLFFENYEHNTSVVKKIHIEVVDKANDMA 155

RESULT 4
US-09-997-579-44
; Sequence 44, Application US/09997579
; Patent No. US20020113203A1
; GENERAL INFORMATION:
; APPLICANT: Cambridge University Technical Services
; TITLE OF INVENTION: A novel family of beta sub-unit proteins from a voltage gated soc
; TITLE OF INVENTION: channel
; FILE REFERENCE: nucleic acids encoding them and therapeutic or diagnostic uses
; FILE REFERENCE: 674558-2001
; CURRENT APPLICATION NUMBER: US/09/997,579
; CURRENT FILING DATE: 2002-04-05
; PRIOR APPLICATION NUMBER: PCT/EP00/01783
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: 60,129,473
; PRIOR FILING DATE: 2000-02-24
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 44
; LENGTH: 218
; TYPE: PRT
; ORGANISM: Rat
US-09-997-579-44

Query Match 52.0%; Score 751; DB 9; Length 218;
Best Local Similarity 92.9%; Pred. No. 1e-65;
Matches 144; Conservative 2; Mismatches 9; Indels 0; Gaps 0;

Qy 1 MGRLLALVVGAAVLSVSSACGGCVVDSETEAVYGMTFKILCISCKRSETNAETFTWTFR 60
Db 1 MGRLLALVVGAAVLSVSSACGGCVVDSETEAVYGMTFKILCISCKRSETNAETFTWTFR 60

Qy 61 QKGTTEFVKILRYENEVLQLEDEDERFEGRVVWNGSRGTDQLDLSIFITNTVYHSGDYE 120
Db 61 QKGTTEFVKILRYENEVLQLEDEDERFEGRVVWNGSRGTDQLDLSIFITNTVYHSGDYE 120

Qy 121 CHVYRLFFENYEHNTSVVKKIHIEVVDKESGAA 155
Db 121 CHVYRLFFENYEHNTSVVKKIHIEVVDKANDMA 155

RESULT 5
US-10-029-191-20
; Sequence 20, Application US/10029191
; Publication No. US20020160453A1
; GENERAL INFORMATION:
; APPLICANT: CURTIS, Rory A.J.
; TITLE OF INVENTION: NOVEL GENE ENCODING A SODIUM CHANNEL BETA-3 SUBUNIT
; FILE REFERENCE: PROTEIN
; FILE REFERENCE: 210147.00XX/SU1
; CURRENT APPLICATION NUMBER: US/10/029,191
; CURRENT FILING DATE: 2001-12-20
; PRIOR APPLICATION NUMBER: 09/569,978

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; PRIOR FILING DATE: 2000-05-12
; PRIOR APPLICATION NUMBER: US 60/134,198
; PRIOR FILING DATE: 1999-05-14
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 20
; LENGTH: 218
; TYPE: PRT
; ORGANISM: Lepus Sp.
US-10-029-191-20

Query Match
  51.9%; Score 749; DB 13; Length 218;
Best Local Similarity 92.9%; Pred. No. 1.6e-65;
Matches 144; Conservative 2; Mismatches 9; Indels 0; Gaps 0;

QY 1 MGRLLALVGAALVSSACGCGVEVDSEAVYGMTFKILCISCKRSETNAETFTETWFR 60
DB 1 MGRLLAFVGAALVSSACGCGVEVDSEAVYGMTFKILCISCKRSETTAETFTWFR 60

QY 61 QKGTBEFVKILRYENEVLQLEEDERFEGRVVWNGSGTKDQLDLSIFITNTVYHSGDYE 120
DB 61 QKGTBEFVKILRYENEVLQLEEDERFEGRVVWNGSGTKDQLDLSIFITNTVYHSGDYQ 120

QY 121 CHVYRLLPENYEHTSVVKKIHIEVDKGESGAA 155
DB 121 CHVYRLLPENYEHTSVVKKIHIEVDKXNRDMA 155

RESULT 6
US-09-997-579-22
; Sequence 22, Application US/09997579
; Patent No. US20020113203A1
; GENERAL INFORMATION:
; APPLICANT: Cambridge University Technical Services
; TITLE OF INVENTION: A novel family of beta sub-unit proteins from a voltage gated so
; TITLE OF INVENTION: channel
; TITLE OF INVENTION: nucleic acids encoding them and therapeutic or diagnostic uses
; FILE REFERENCE: 674558-2001
; CURRENT APPLICATION NUMBER: US/09/997,579
; PRIOR FILING DATE: 2002-04-05
; PRIOR APPLICATION NUMBER: PCT/EP00/01783
; PRIOR FILING DATE: 2000-02-24
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 22
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-997-579-22

Query Match
  17.9%; Score 258.5; DB 9; Length 159;
Best Local Similarity 43.2%; Pred. No. 3.1e-17;
Matches 64; Conservative 20; Mismatches 57; Indels 7; Gaps 4;

QY 5 LALVGAALVSSACGCGVEVDSEAVYGMTFKILCISCKRSETNAETFTETWFRQKGT 64
DB 10 LASLVLIYVWVSCFPVCEVPSETEAVQGNPKMLRCISCKRSEETVVEWFFRPEGG 69

QY 65 EEFVKILRYENEVLQLEEDERFEGRVVWNGSGTKDQLDLSIFITNTVYHSGDYCHVY 124
DB 70 KDFL-IYEYRNHGHQVEESP--FQGRLOWNGS---KDLQDVSITVLNLTNDSGLYTCNVS 123

QY 125 RLFFENYEHTSVVKKIHIEVDK-GE 151
DB 124 REFEFAHRPFVKTTLPIRVTERAGE 151

RESULT 7
US-09-997-579-22
; Sequence 22, Application US/09997579
; Patent No. US20020113203A1
; GENERAL INFORMATION:
; APPLICANT: Cambridge University Technical Services
; TITLE OF INVENTION: A novel family of beta sub-unit proteins from a voltage gated so
; TITLE OF INVENTION: channel
; TITLE OF INVENTION: nucleic acids encoding them and therapeutic or diagnostic uses
; FILE REFERENCE: 674558-2001
; CURRENT APPLICATION NUMBER: US/09/997,579
; PRIOR FILING DATE: 2002-04-05
; PRIOR APPLICATION NUMBER: PCT/EP00/01783
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: 60,129,473
; PRIOR FILING DATE: 2000-02-24
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 22
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-997-579-22

Query Match
  17.9%; Score 258.5; DB 9; Length 159;
Best Local Similarity 43.2%; Pred. No. 3.1e-17;
Matches 64; Conservative 20; Mismatches 57; Indels 7; Gaps 4;

QY 5 LALVGAALVSSACGCGVEVDSEAVYGMTFKILCISCKRSETNAETFTETWFRQKGT 64
DB 10 LASLVLIYVWVSCFPVCEVPSETEAVQGNPKMLRCISCKRSEETVVEWFFRPEGG 69

QY 65 EEFVKILRYENEVLQLEEDERFEGRVVWNGSGTKDQLDLSIFITNTVYHSGDYCHVY 124
DB 70 KDFL-IYEYRNHGHQVEESP--FQGRLOWNGS---KDLQDVSITVLNLTNDSGLYTCNVS 123

QY 125 RLFFENYEHTSVVKKIHIEVDK-GE 151
DB 124 REFEFAHRPFVKTTLPIRVTERAGE 151

RESULT 8
US-10-142-201B-11
; Sequence 11, Application US/10142201B
; Publication No. US20030022205A1
; GENERAL INFORMATION:
; APPLICANT: Millennium Pharmaceuticals Inc.
; APPLICANT: Curtis, Rory A.J.
; TITLE OF INVENTION: 98359, A SODIUM CHANNEL BETA 4 SUBUNIT,
; TITLE OF INVENTION: AND USES THEREFOR
; FILE REFERENCE: MP12001-108PIRN(M)
; CURRENT APPLICATION NUMBER: US/10/142,201B
; CURRENT FILING DATE: 2002-05-09
; PRIOR APPLICATION NUMBER: US 60/289,893
; PRIOR FILING DATE: 2001-05-09
; NUMBER OF SEQ ID NOS: 12
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 11
; LENGTH: 215
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-142-201B-11

Query Match
  17.9%; Score 258.5; DB 14; Length 215;
Best Local Similarity 43.2%; Pred. No. 4.6e-17;
Matches 64; Conservative 20; Mismatches 57; Indels 7; Gaps 4;

QY 5 LALVGAALVSSACGCGVEVDSEAVYGMTFKILCISCKRSETNAETFTETWFRQKGT 64
DB 10 LASLVLIYVWVSCFPVCEVPSETEAVQGNPKMLRCISCKRSEETVVEWFFRPEGG 69

QY 65 EEFVKILRYENEVLQLEEDERFEGRVVWNGSGTKDQLDLSIFITNTVYHSGDYCHVY 124
DB 70 KDFL-IYEYRNHGHQVEESP--FQGRLOWNGS---KDLQDVSITVLNLTNDSGLYTCNVS 123

QY 125 RLFFENYEHTSVVKKIHIEVDK-GE 151
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; PRIOR FILING DATE: 2000-05-12
; PRIOR APPLICATION NUMBER: US 60/134,198
; PRIOR FILING DATE: 1999-05-14
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 20
; LENGTH: 218
; TYPE: PRT
; ORGANISM: Lepus Sp.
US-10-029-191-20

Query Match
  51.9%; Score 749; DB 13; Length 218;
Best Local Similarity 92.9%; Pred. No. 1.6e-65;
Matches 144; Conservative 2; Mismatches 9; Indels 0; Gaps 0;

QY 1 MGRLLALVGAALVSSACGCGVEVDSEAVYGMTFKILCISCKRSETNAETFTETWFR 60
DB 1 MGRLLAFVGAALVSSACGCGVEVDSEAVYGMTFKILCISCKRSETTAETFTWFR 60

QY 61 QKGTBEFVKILRYENEVLQLEEDERFEGRVVWNGSGTKDQLDLSIFITNTVYHSGDYE 120
DB 61 QKGTBEFVKILRYENEVLQLEEDERFEGRVVWNGSGTKDQLDLSIFITNTVYHSGDYQ 120

QY 121 CHVYRLLPENYEHTSVVKKIHIEVDKGESGAA 155
DB 121 CHVYRLLPENYEHTSVVKKIHIEVDKXNRDMA 155

RESULT 6
US-09-997-579-22
; Sequence 22, Application US/09997579
; Patent No. US20020113203A1
; GENERAL INFORMATION:
; APPLICANT: Cambridge University Technical Services
; TITLE OF INVENTION: A novel family of beta sub-unit proteins from a voltage gated so
; TITLE OF INVENTION: channel
; TITLE OF INVENTION: nucleic acids encoding them and therapeutic or diagnostic uses
; FILE REFERENCE: 674558-2001
; CURRENT APPLICATION NUMBER: US/09/997,579
; PRIOR FILING DATE: 2002-04-05
; PRIOR APPLICATION NUMBER: PCT/EP00/01783
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: 60,129,473
; PRIOR FILING DATE: 2000-02-24
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 22
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-997-579-22

Query Match
  17.9%; Score 258.5; DB 9; Length 159;
Best Local Similarity 43.2%; Pred. No. 3.1e-17;
Matches 64; Conservative 20; Mismatches 57; Indels 7; Gaps 4;

QY 5 LALVGAALVSSACGCGVEVDSEAVYGMTFKILCISCKRSETNAETFTETWFRQKGT 64
DB 10 LASLVLIYVWVSCFPVCEVPSETEAVQGNPKMLRCISCKRSEETVVEWFFRPEGG 69

QY 65 EEFVKILRYENEVLQLEEDERFEGRVVWNGSGTKDQLDLSIFITNTVYHSGDYCHVY 124
DB 70 KDFL-IYEYRNHGHQVEESP--FQGRLOWNGS---KDLQDVSITVLNLTNDSGLYTCNVS 123

QY 125 RLFFENYEHTSVVKKIHIEVDK-GE 151
DB 124 REFEFAHRPFVKTTLPIRVTERAGE 151

RESULT 7
US-09-997-579-22
; Sequence 22, Application US/09997579
; Patent No. US20020113203A1
; GENERAL INFORMATION:
; APPLICANT: Cambridge University Technical Services
; TITLE OF INVENTION: A novel family of beta sub-unit proteins from a voltage gated so
; TITLE OF INVENTION: channel
; TITLE OF INVENTION: nucleic acids encoding them and therapeutic or diagnostic uses
; FILE REFERENCE: 674558-2001
; CURRENT APPLICATION NUMBER: US/09/997,579
; PRIOR FILING DATE: 2002-04-05
; PRIOR APPLICATION NUMBER: PCT/EP00/01783
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: 60,129,473
; PRIOR FILING DATE: 2000-02-24
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 22
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-997-579-22

Query Match
  17.9%; Score 258.5; DB 9; Length 159;
Best Local Similarity 43.2%; Pred. No. 3.1e-17;
Matches 64; Conservative 20; Mismatches 57; Indels 7; Gaps 4;

QY 5 LALVGAALVSSACGCGVEVDSEAVYGMTFKILCISCKRSETNAETFTETWFRQKGT 64
DB 10 LASLVLIYVWVSCFPVCEVPSETEAVQGNPKMLRCISCKRSEETVVEWFFRPEGG 69

QY 65 EEFVKILRYENEVLQLEEDERFEGRVVWNGSGTKDQLDLSIFITNTVYHSGDYCHVY 124
DB 70 KDFL-IYEYRNHGHQVEESP--FQGRLOWNGS---KDLQDVSITVLNLTNDSGLYTCNVS 123

QY 125 RLFFENYEHTSVVKKIHIEVDK-GE 151
DB 124 REFEFAHRPFVKTTLPIRVTERAGE 151

RESULT 8
US-10-142-201B-11
; Sequence 11, Application US/10142201B
; Publication No. US20030022205A1
; GENERAL INFORMATION:
; APPLICANT: Millennium Pharmaceuticals Inc.
; APPLICANT: Curtis, Rory A.J.
; TITLE OF INVENTION: 98359, A SODIUM CHANNEL BETA 4 SUBUNIT,
; TITLE OF INVENTION: AND USES THEREFOR
; FILE REFERENCE: MP12001-108PIRN(M)
; CURRENT APPLICATION NUMBER: US/10/142,201B
; CURRENT FILING DATE: 2002-05-09
; PRIOR APPLICATION NUMBER: US 60/289,893
; PRIOR FILING DATE: 2001-05-09
; NUMBER OF SEQ ID NOS: 12
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 11
; LENGTH: 215
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-142-201B-11

Query Match
  17.9%; Score 258.5; DB 14; Length 215;
Best Local Similarity 43.2%; Pred. No. 4.6e-17;
Matches 64; Conservative 20; Mismatches 57; Indels 7; Gaps 4;

QY 5 LALVGAALVSSACGCGVEVDSEAVYGMTFKILCISCKRSETNAETFTETWFRQKGT 64
DB 10 LASLVLIYVWVSCFPVCEVPSETEAVQGNPKMLRCISCKRSEETVVEWFFRPEGG 69

QY 65 EEFVKILRYENEVLQLEEDERFEGRVVWNGSGTKDQLDLSIFITNTVYHSGDYCHVY 124
DB 70 KDFL-IYEYRNHGHQVEESP--FQGRLOWNGS---KDLQDVSITVLNLTNDSGLYTCNVS 123

QY 125 RLFFENYEHTSVVKKIHIEVDK-GE 151
```


GenCore version 5.1.6
Copyright (c) 1993 - 2004 CompuGen Ltd.

OM protein - protein search, using sw model

Run on: September 21, 2004, 16:59:29 ; Search time 33 Seconds
(without alignments)
419.265 Million cell updates/sec

Title: US-09-875-456A-14
Perfect score: 1444
Sequence: 1 MGRLLALVGAALVSSACGG.....QRVLGLPGIIIRCVSEGVV 268

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 389414 seqs, 51625971 residues

Total number of hits satisfying chosen parameters: 389414

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued Patents_AA*

- 1: /cgn2_6/ptodata/2/iaa/5A_COMB.pap.*
- 2: /cgn2_6/ptodata/2/iaa/5B_COMB.pap.*
- 3: /cgn2_6/ptodata/2/iaa/6A_COMB.pap.*
- 4: /cgn2_6/ptodata/2/iaa/6B_COMB.pap.*
- 5: /cgn2_6/ptodata/2/iaa/6CTUS_COMB.pap.*
- 6: /cgn2_6/ptodata/2/iaa/backfiles1.pap.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Query Match	Score	Length	DB ID	Description
1	751	52.0	218	4	US-09-997-579-44
2	258.5	17.9	159	4	US-09-997-579-22
3	258.5	17.9	215	4	US-09-997-579-2
4	257.5	17.8	159	4	US-09-997-579-23
5	257.5	17.8	215	4	US-09-997-579-1
6	115.5	8.0	209	4	US-09-430-503-20
7	115.5	8.0	209	4	US-09-430-503-24
8	115.5	8.0	269	4	US-09-430-503-4
9	115.5	8.0	269	4	US-09-430-503-6
10	115.5	8.0	269	4	US-09-430-503-8
11	114.5	7.9	159	4	US-09-430-503-34
12	114.5	7.9	159	4	US-09-430-503-38
13	114.5	7.9	199	4	US-09-430-503-42
14	114.5	7.9	199	4	US-09-430-503-46
15	114.5	7.9	209	4	US-09-430-503-18
16	114.5	7.9	209	4	US-09-430-503-22
17	114.5	7.9	269	4	US-09-430-503-2
18	112.5	7.8	159	4	US-09-430-503-36
19	112.5	7.8	159	4	US-09-430-503-40
20	112.5	7.8	199	4	US-09-430-503-44
21	112.5	7.8	199	4	US-09-430-503-48
22	107.5	7.4	270	4	US-09-430-503-26
23	107.5	7.4	270	4	US-09-430-503-28
24	107.5	7.4	270	4	US-09-430-503-30
25	107.5	7.4	270	4	US-09-430-503-32
26	98	6.8	380	3	US-08-459-953A-9
27	98	6.8	380	4	US-09-393-212-9

28	94	6.5	358	1	US-08-176-620A-12	Sequence 12, Appl
29	94	6.5	358	2	US-08-461-985-12	Sequence 12, Appl
30	94	6.5	358	4	US-09-457-040B-4	Sequence 4, Appl
31	94	6.5	360	4	US-09-457-040B-3	Sequence 3, Appl
32	94	6.5	364	4	US-09-457-040B-34	Sequence 34, Appl
33	94	6.5	415	1	US-08-176-620A-4	Sequence 4, Appl
34	94	6.5	415	1	US-08-463-862-4	Sequence 4, Appl
35	94	6.5	415	2	US-08-461-985-4	Sequence 4, Appl
36	94	6.5	415	2	US-08-458-887-4	Sequence 4, Appl
37	94	6.5	415	3	US-08-932-787B-4	Sequence 4, Appl
38	94	6.5	415	3	US-08-932-012C-4	Sequence 4, Appl
39	94	6.5	415	4	US-08-888-818C-4	Sequence 4, Appl
40	94	6.5	604	4	US-09-417-197-59	Sequence 59, Appl
41	94	6.5	605	4	US-09-417-197-41	Sequence 41, Appl
42	93	6.4	246	4	US-09-252-991A-23594	Sequence 23594, A
43	91.5	6.3	292	4	US-09-205-258-1116	Sequence 1116, Ap
44	91.5	6.3	4302	4	US-09-052-469-8	Sequence 8, Appl
45	91.5	6.3	4302	4	US-08-422-582-8	Sequence 8, Appl

ALIGNMENTS

RESULT 1
US-09-997-579-44
; Sequence 44, Application US/09997579
; Patent No. 6593565
; GENERAL INFORMATION:
; APPLICANT: Cambridge University Technical Services
; TITLE OF INVENTION: A novel family of beta sub-unit proteins from a voltage gated so
; TITLE OF INVENTION: channel
; TITLE OF INVENTION: nucleic acids encoding them and therapeutic or diagnostic uses
; FILE REFERENCE: 674558-2001
; CURRENT APPLICATION NUMBER: US/09/997,579
; CURRENT FILING DATE: 2002-04-05
; PRIOR APPLICATION NUMBER: PCT/EP00/01783
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: 60,129,473
; PRIOR FILING DATE: 2000-02-24
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 44
; LENGTH: 218
; TYPE: PRT
; ORGANISM: Rat
US-09-997-579-44

Query Match	52.0%;	Score 751;	DB 4;	Length 218;
Best Local Similarity	92.9%;	Pred. No. 1.6e-72;		
Matches 144;	Conservative 2;	Mismatches 9;	Indels 0;	Gaps 0;
Qy	1	MGRLLALVGAALVSSACGGCVSDSEAVYGMFKILCISCKRRSETNAETFTWTFR	60	
Db	1	MGRLLALVGAALVSSACGGCVSDSEAVYGMFKILCISCKRRSETNAETFTWTFR	60	
Qy	61	QKGTSEFVKILYENVEVLQLEDERPEGRVWNGSRGKDLQLSIFITVNTYHSGDYE	120	
Db	61	QKGTSEFVKILYENVEVLQLEDERPEGRVWNGSRGKDLQLSIFITVNTYHSGDYE	120	
Qy	121	CHVYRLFFPNYEHNTSVVKKIHIEVDKSGGAA	155	
Db	121	CHVYRLFFPNYEHNTSVVKKIHIEVDKSGGAA	155	

RESULT 2
US-09-997-579-22
; Sequence 22, Application US/09997579
; Patent No. 6593565
; GENERAL INFORMATION:
; APPLICANT: Cambridge University Technical Services
; TITLE OF INVENTION: A novel family of beta sub-unit proteins from a voltage gated so
; TITLE OF INVENTION: channel
; TITLE OF INVENTION: nucleic acids encoding them and therapeutic or diagnostic uses

```
; FILE REFERENCE: 674558-2001
; CURRENT APPLICATION NUMBER: US/09/997,579
; CURRENT FILING DATE: 2002-04-05
; PRIOR APPLICATION NUMBER: PCT/EP00/01783
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: 60,129,473
; PRIOR FILING DATE: 2000-02-24
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 22
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-997-579-22

Query Match          17.9%; Score 258.5; DB 4; Length 159;
Best Local Similarity 43.2%; Pred. No. 8.2e-20;
Matches 64; Conservative 20; Mismatches 57; Indels 7; Gaps 4;

QY 5 LALVGAALVSSACGCGVEVDSEAVYGMTFKILCISKRRSETNAETFTWFRQKGT 64
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 10 LASLVLIYVSVCFVCPVCEVPESEAVQGNPKLRCISCMKREEVEATTVVEWFRPEG 69
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 65 EEFVKILRYENEVLQLEDEDEREGRVVWNGSGTKDLODLSIFITNTVYHSGDYEGHY 124
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 70 KDPL-IYEYRNHGHQVEESP--FQGRLOWNGS---KDQDVSITVLNVTLNDSGLYTCNV 123
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 125 RLLFFENYEHTSVVVKIHIEVDK-GE 151
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 124 REFEEAHRPFVKTTRLIPLRVTEAGE 151
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
```

```
RESULT 3
US-09-997-579-2
; Sequence 2, Application US/09997579
; Patent No. 6593565
; GENERAL INFORMATION:
; APPLICANT: Cambridge University Technical Services
; TITLE OF INVENTION: A novel family of beta sub-unit proteins from a voltage gated soc
; TITLE OF INVENTION: channel
; FILE REFERENCE: 674558-2001
; CURRENT APPLICATION NUMBER: US/09/997,579
; CURRENT FILING DATE: 2002-04-05
; PRIOR APPLICATION NUMBER: PCT/EP00/01783
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: 60,129,473
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 2
; LENGTH: 215
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-997-579-2

Query Match          17.9%; Score 258.5; DB 4; Length 215;
Best Local Similarity 43.2%; Pred. No. 1.3e-19;
Matches 64; Conservative 20; Mismatches 57; Indels 7; Gaps 4;

QY 5 LALVGAALVSSACGCGVEVDSEAVYGMTFKILCISKRRSETNAETFTWFRQKGT 64
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 10 LASLVLIYVSVCFVCPVCEVPESEAVQGNPKLRCISCMKREEVEATTVVEWFRPEG 69
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 65 EEFVKILRYENEVLQLEDEDEREGRVVWNGSGTKDLODLSIFITNTVYHSGDYEGHY 124
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 70 KDPL-IYEYRNHGHQVEESP--FQGRLOWNGS---KDQDVSITVLNVTLNDSGLYTCNV 123
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 125 RLLFFENYEHTSVVVKIHIEVDK-GE 151
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 124 REFEEAHRPFVKTTRLIPLRVTEAGE 151
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
```

```
RESULT 4
US-09-997-579-23
; Sequence 23, Application US/09997579
; Patent No. 6593565
; GENERAL INFORMATION:
; APPLICANT: Cambridge University Technical Services
; TITLE OF INVENTION: A novel family of beta sub-unit proteins from a voltage gated soc
; TITLE OF INVENTION: channel
; FILE REFERENCE: 674558-2001
; CURRENT APPLICATION NUMBER: US/09/997,579
; CURRENT FILING DATE: 2002-04-05
; PRIOR APPLICATION NUMBER: PCT/EP00/01783
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: 60,129,473
; PRIOR FILING DATE: 2000-02-24
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 23
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Rat
US-09-997-579-23

Query Match          17.8%; Score 257.5; DB 4; Length 159;
Best Local Similarity 44.7%; Pred. No. 1e-19;
Matches 59; Conservative 19; Mismatches 47; Indels 7; Gaps 4;

QY 21 CVEVDSEAVYGMTFKILCISKRRSETNAETFTWFRQKGTBEFVKILRYENEVLQL 80
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 26 CVEVPESEAVQGNPKLRCISCMKREEVEATTVVEWFRPEGKDFL-IYEYRNHGHQEV 84
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 81 EDERPEGRVWNGSGTKDLODLSIFITNTVYHSGDYEGHYVLLFFENYEHTSVVK 140
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 85 ESP--FQGRLOWNGS---KDQDVSITVLNVTLNDSGLYTCNVSRPEFEAHRPFVKTTR 139
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 141 KIHIEVDK-GE 151
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 140 LIPLRVTEAGE 151
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
```

```
RESULT 5
US-09-997-579-1
; Sequence 1, Application US/09997579
; Patent No. 6593565
; GENERAL INFORMATION:
; APPLICANT: Cambridge University Technical Services
; TITLE OF INVENTION: A novel family of beta sub-unit proteins from a voltage gated soc
; TITLE OF INVENTION: channel
; FILE REFERENCE: 674558-2001
; CURRENT APPLICATION NUMBER: US/09/997,579
; CURRENT FILING DATE: 2002-04-05
; PRIOR APPLICATION NUMBER: PCT/EP00/01783
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: 60,129,473
; PRIOR FILING DATE: 2000-02-24
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 1
; LENGTH: 215
; TYPE: PRT
; ORGANISM: Rat
US-09-997-579-1

Query Match          17.8%; Score 257.5; DB 4; Length 215;
Best Local Similarity 44.7%; Pred. No. 1.6e-19;
Matches 59; Conservative 19; Mismatches 47; Indels 7; Gaps 4;

QY 21 CVEVDSEAVYGMTFKILCISKRRSETNAETFTWFRQKGTBEFVKILRYENEVLQL 80
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 26 CVEVPESEAVQGNPKLRCISCMKREEVEATTVVEWFRPEGKDFL-IYEYRNHGHQEV 84
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
```



```
Db 77 QPEGADTTVSFFHYSQGVYLGNYPPFKDRISW---AGDLKKDASININENQFIHNGTY 133
QY 120 ECHV 123
Db 134 ICDV 137

RESULT 10
US-09-430-503-8
; Sequence 8, Application US/09430503
; Patent No. 6355786
; GENERAL INFORMATION:
; APPLICANT: Zhao, Zhizhuang
; TITLE OF INVENTION: PURIFIED AND ISOLATED PROTEIN ZERO RELATED (PZR) AND
; TITLE OF INVENTION: THERAPEUTIC AND SCREENING METHODS USING SAME
; FILE REFERENCE: Attorney Docket No. 6355786 1242-11/2
; CURRENT APPLICATION NUMBER: US/09/430,503
; CURRENT FILING DATE: 1999-10-29
; NUMBER OF SEQ ID NOS: 49
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 8
; LENGTH: 269
; TYPE: PRT
; ORGANISM: Homo sapiens
; US-09-430-503-8

Query Match 8.0%; Score 115.5; DB 4; Length 269;
Best Local Similarity 30.6%; Pred. No. 0.0004;
Matches 38; Conservative 17; Mismatches 62; Indels 7; Gaps 4;
QY 3 RLALVVGAAAL-VSSAGCGVEVDSETE--AVYGMTFKILCISKRRSETNAETFTWTF 59
Db 18 RMLWSVLAALGGLTAGVSALEVYTPKEIFVANGTQGLTC-KFKSTSTTGGTGSVWSF 76
QY 60 ROKGTEEFVKILRYENEVLQLEEDERFEGRVVWNGSRGTKQLDLSIFITNTVYHNSGDY 119
Db 77 QPEGADTTVSFFHYSQGVYLGNYPPFKDRISW---AGDLKKDASININENQFIHNGTY 133
QY 120 ECHV 123
Db 134 ICDV 137

RESULT 11
US-09-430-503-34
; Sequence 34, Application US/09430503
; Patent No. 6355786
; GENERAL INFORMATION:
; APPLICANT: Zhao, Zhizhuang
; TITLE OF INVENTION: PURIFIED AND ISOLATED PROTEIN ZERO RELATED (PZR) AND
; TITLE OF INVENTION: THERAPEUTIC AND SCREENING METHODS USING SAME
; FILE REFERENCE: Attorney Docket No. 6355786 1242-11/2
; CURRENT APPLICATION NUMBER: US/09/430,503
; CURRENT FILING DATE: 1999-10-29
; NUMBER OF SEQ ID NOS: 49
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 34
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Homo sapiens
; US-09-430-503-34

Query Match 7.9%; Score 114.5; DB 4; Length 159;
Best Local Similarity 30.6%; Pred. No. 0.00024;
Matches 38; Conservative 15; Mismatches 64; Indels 7; Gaps 4;
QY 3 RLALVVGAAALVSSAGC-GCVEVDSETE--AVYGMTFKILCISKRRSETNAETFTWTF 59
Db 18 RMLWSVLAALGGLTAGVSALEVYTPKEIFVANGTQGLTC-KFKSTSTTGGTGSVWSF 76
QY 60 ROKGTEEFVKILRYENEVLQLEEDERFEGRVVWNGSRGTKQLDLSIFITNTVYHNSGDY 119
Db 77 QPEGADTTVSFFHYSQGVYLGNYPPFKDRISW---AGDLKKDASININENQFIHNGTY 133
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QY 120 ECHV 123
Db 134 ICDV 137
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```
RESULT 12
US-09-430-503-38
; Sequence 38, Application US/09430503
; Patent No. 6355786
; GENERAL INFORMATION:
; APPLICANT: Zhao, Zhizhuang
; TITLE OF INVENTION: PURIFIED AND ISOLATED PROTEIN ZERO RELATED (PZR) AND
; TITLE OF INVENTION: THERAPEUTIC AND SCREENING METHODS USING SAME
; FILE REFERENCE: Attorney Docket No. 6355786 1242-11/2
; CURRENT APPLICATION NUMBER: US/09/430,503
; CURRENT FILING DATE: 1999-10-29
; NUMBER OF SEQ ID NOS: 49
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 38
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Homo sapiens
; US-09-430-503-38
```

```
Query Match 7.9%; Score 114.5; DB 4; Length 159;
Best Local Similarity 30.6%; Pred. No. 0.00024;
Matches 38; Conservative 15; Mismatches 64; Indels 7; Gaps 4;
QY 3 RLALVVGAAALVSSAGC-GCVEVDSETE--AVYGMTFKILCISKRRSETNAETFTWTF 59
Db 18 RMLWSVLAALGGLTAGVSALEVYTPKEIFVANGTQGLTC-KFKSTSTTGGTGSVWSF 76
QY 60 ROKGTEEFVKILRYENEVLQLEEDERFEGRVVWNGSRGTKQLDLSIFITNTVYHNSGDY 119
Db 77 QPEGADTTVSFFHYSQGVYLGNYPPFKDRISW---AGDLKKDASININENQFIHNGTY 133
QY 120 ECHV 123
Db 134 ICDV 137
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RESULT 13
US-09-430-503-42
; Sequence 42, Application US/09430503
; Patent No. 6355786
; GENERAL INFORMATION:
; APPLICANT: Zhao, Zhizhuang
; TITLE OF INVENTION: PURIFIED AND ISOLATED PROTEIN ZERO RELATED (PZR) AND
; TITLE OF INVENTION: THERAPEUTIC AND SCREENING METHODS USING SAME
; FILE REFERENCE: Attorney Docket No. 6355786 1242-11/2
; CURRENT APPLICATION NUMBER: US/09/430,503
; CURRENT FILING DATE: 1999-10-29
; NUMBER OF SEQ ID NOS: 49
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 42
; LENGTH: 199
; TYPE: PRT
; ORGANISM: Homo sapiens
; US-09-430-503-42
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```
Query Match 7.9%; Score 114.5; DB 4; Length 199;
Best Local Similarity 30.6%; Pred. No. 0.0003;
Matches 38; Conservative 15; Mismatches 64; Indels 7; Gaps 4;
QY 3 RLALVVGAAALVSSAGC-GCVEVDSETE--AVYGMTFKILCISKRRSETNAETFTWTF 59
Db 18 RMLWSVLAALGGLTAGVSALEVYTPKEIFVANGTQGLTC-KFKSTSTTGGTGSVWSF 76
QY 60 ROKGTEEFVKILRYENEVLQLEEDERFEGRVVWNGSRGTKQLDLSIFITNTVYHNSGDY 119
Db 77 QPEGADTTVSFFHYSQGVYLGNYPPFKDRISW---AGDLKKDASININENQFIHNGTY 133
```

QY 120 ECHV 123
Db 134 ICDV 137

RESULT 14

US-09-430-503-46

; Sequence 46, Application US/09430503

; Patent No. 6355786

; GENERAL INFORMATION:

; APPLICANT: Zhao, Zhizhuang

; TITLE OF INVENTION: PURIFIED AND ISOLATED PROTEIN ZERO RELATED (PZR) AND

; TITLE OF INVENTION: THERAPEUTIC AND SCREENING METHODS USING SAME

; FILE REFERENCE: Attorney Docket No. 6355786 1242-11/2

; CURRENT APPLICATION NUMBER: US/09/430,503

; CURRENT FILING DATE: 1999-10-29

; NUMBER OF SEQ ID NOS: 49

; SOFTWARE: PatentIn Ver. 2.0

; SEQ ID NO 46

; LENGTH: 199

; TYPE: PRT

; ORGANISM: Homo sapiens

US-09-430-503-46

Query Match 7.9%; Score 114.5; DB 4; Length 199;

Best Local Similarity 30.6%; Pred. No. 0.00033;

Matches 38; Conservative 15; Mismatches 64; Indels 7; Gaps 4;

QY 3 RLLALVGAALVSSACG-GCVEVDSETE--AVYGMTFKILCISCKRRSETNAETFTETWF 59

Db 18 RLMVSLAALGLLTAGVSALEVTYKEIFVANGTQGLTC-KPKSTSTTGGTSLVSWSP 76

QY 60 ROKGTEEFVKILRYENEVLQLEEDERFEGRVVWNGSRGTDLQDLSIFITVYVNHSGDY 119

Db 77 QPEGADTTVSFFHYSGQVYLGNYPPFKDRISW---AGDLKDKDASINENNQFTHNGTY 133

QY 120 ECHV 123
Db 134 ICDV 137

RESULT 15

US-09-430-503-18

; Sequence 18, Application US/09430503

; Patent No. 6355786

; GENERAL INFORMATION:

; APPLICANT: Zhao, Zhizhuang

; TITLE OF INVENTION: PURIFIED AND ISOLATED PROTEIN ZERO RELATED (PZR) AND

; TITLE OF INVENTION: THERAPEUTIC AND SCREENING METHODS USING SAME

; FILE REFERENCE: Attorney Docket No. 6355786 1242-11/2

; CURRENT APPLICATION NUMBER: US/09/430,503

; CURRENT FILING DATE: 1999-10-29

; NUMBER OF SEQ ID NOS: 49

; SOFTWARE: PatentIn Ver. 2.0

; SEQ ID NO 18

; LENGTH: 209

; TYPE: PRT

; ORGANISM: Homo sapiens

US-09-430-503-18

Query Match 7.9%; Score 114.5; DB 4; Length 209;

Best Local Similarity 30.6%; Pred. No. 0.00036;

Matches 38; Conservative 15; Mismatches 64; Indels 7; Gaps 4;

QY 3 RLLALVGAALVSSACG-GCVEVDSETE--AVYGMTFKILCISCKRRSETNAETFTETWF 59

Db 18 RLMVSLAALGLLTAGVSALEVTYKEIFVANGTQGLTC-KPKSTSTTGGTSLVSWSP 76

QY 60 ROKGTEEFVKILRYENEVLQLEEDERFEGRVVWNGSRGTDLQDLSIFITVYVNHSGDY 119

Db 77 QPEGADTTVSFFHYSGQVYLGNYPPFKDRISW---AGDLKDKDASINENNQFTHNGTY 133

QY 120 ECHV 123

Db 134 ICDV 137

Search completed: September 21, 2004, 17:16:54
Job time : 34 secs

